

Sequence listing

<110> SWITCH Biotech AG

<120> Polypeptides or nucleic acids encoding these of a family of G-protein coupled receptors and their use for the diagnosis or treatment of disorders, for example skin disorders and their use for the identification of pharmacologically active substances

<130> S34321US1

<160> 21

<170> WORD6.0, PC-DOS/MS-DOS

<210> 1

<211> 331

<212> PRT

<213> Mus musculus

<400> 1

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Gly Ser Ile Asn Ile Arg Ile Leu Ile Pro Lys Leu Met Ile Ile Ile  
35 40 45

Phe Gly Leu Val Gly Leu Met Gly Asn Ala Ile Val Phe Trp Leu Leu  
50 55 60

Gly Phe His Leu Arg Arg Asn Ala Phe Ser Val Tyr Ile Leu Asn Leu  
65 70 75 80

Ala Leu Ala Asp Phe Leu Phe Leu Ser Ser Ile Ile Ala Ser Thr

85

90

95

Leu Phe Leu Leu Lys Val Ser Tyr Leu Ser Ile Ile Phe His Leu Cys  
100 105 110

Phe Asn Thr Ile Met Met Val Val Tyr Ile Thr Gly Ile Ser Met Leu  
115 120 125

Ser Ala Ile Ser Thr Glu Cys Cys Leu Ser Val Leu Cys Pro Thr Trp  
130 135 140

Tyr Arg Cys His Arg Pro Val His Thr Ser Thr Val Met Cys Ala Val  
145 150 155 160

Ile Trp Val Leu Ser Leu Leu Ile Cys Ile Leu Asn Ser Tyr Phe Cys  
165 170 175

Ala Val Leu His Thr Arg Tyr Asp Asn Asp Asn Glu Cys Leu Ala Thr  
180 185 190

Asn Ile Phe Thr Ala Ser Tyr Met Ile Phe Leu Leu Val Val Leu Cys  
195 200 205

Leu Ser Ser Leu Ala Leu Leu Ala Arg Leu Phe Cys Gly Ala Gly Gln  
210 215 220

Met Lys Leu Thr Arg Phe His Val Thr Ile Leu Leu Thr Leu Leu Val  
225 230 235 240

Phe Leu Leu Cys Gly Leu Pro Phe Val Ile Tyr Cys Ile Leu Leu Phe  
245 250 255

Lys Ile Lys Asp Asp Phe His Val Leu Asp Val Asn Leu Tyr Leu Ala  
260 265 270

Leu Glu Val Leu Thr Ala Ile Asn Ser Cys Ala Asn Pro Ile Ile Tyr

275

280

285

Phe Phe Val Gly Ser Phe Arg His Gln Leu Lys His Gln Thr Leu Lys  
290 295 300

Met Val Leu Gln Ser Ala Leu Gln Asp Thr Pro Glu Thr Ala Glu Asn  
305 310 315 320

Met Val Glu Met Ser Ser Asn Lys Ala Glu Pro

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100

<210> 2

<211> 321

<212> PRT

<213> Homo sapiens

<400> 2

Met Asn Gln Thr Leu Asn Ser Ser Gly Thr Val Glu Ser Ala Leu Asn  
1 5 10 15

Tyr Ser Arg Gly Ser Thr Val His Thr Ala Tyr Leu Val Leu Ser Ser  
20 25 30

Leu Ala Met Phe Thr Cys Leu Cys Gly Met Ala Gly Asn Ser Met Val  
35 40 45

Ile Trp Leu Leu Gly Phe Arg Met His Arg Asn Pro Phe Cys Ile Tyr  
50 55 60

Ile Leu Asn Leu Ala Ala Ala Asp Leu Leu Phe Leu Phe Ser Met Ala  
65 70 75 80

Ser Thr Leu Ser Leu Glu Thr Gln Pro Leu Val Asn Thr Thr Asp Lys  
85 90 95

Val His Glu Leu Met Lys Arg Leu Met Tyr Phe Ala Tyr Thr Val Gly

100

105

110

Leu Ser Leu Leu Thr Ala Ile Ser Thr Gln Arg Cys Leu Ser Val Leu

115

120

125

Phe Pro Ile Trp Phe Lys Cys His Arg Pro Arg His Leu Ser Ala Trp

130

135

140

Val Cys Gly Leu Leu Trp Thr Leu Cys Leu Leu Met Asn Gly Leu Thr

145

150

155

160

Ser Ser Phe Cys Ser Lys Phe Leu Lys Phe Asn Glu Asp Arg Cys Phe

165

170

175

Arg Val Asp Met Val Gln Ala Ala Leu Ile Met Gly Val Leu Thr Pro

180

185

190

Val Met Thr Leu Ser Ser Leu Thr Leu Phe Val Trp Val Arg Arg Ser

195

200

205

Ser Gln Gln Trp Arg Arg Gln Pro Thr Arg Leu Phe Val Val Val Leu

210

215

220

Ala Ser Val Leu Val Phe Leu Ile Cys Ser Leu Pro Leu Ser Ile Tyr

225

230

235

240

Trp Phe Val Leu Tyr Trp Leu Ser Leu Pro Pro Glu Met Gln Val Leu

245

250

255

Cys Phe Ser Leu Ser Arg Leu Ser Ser Ser Val Ser Ser Ala Asn

260

265

270

Pro Val Ile Tyr Phe Leu Val Gly Ser Arg Arg Ser His Arg Leu Pro

275

280

285

Thr Arg Ser Leu Gly Thr Val Leu Gln Gln Ala Leu Arg Glu Glu Pro

290

295

300

Glu Leu Glu Gly Gly Glu Thr Pro Thr Val Gly Thr Asn Glu Met Gly

305

310

315

320

Ala

<210> 3

<211> 325

<212> PRT

<213> Mus musculus

<400> 3

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1

5

10

15

Gly Ser Ser Tyr Thr Asn Ser Val Asp Cys Phe Phe Lys Ile Gln Val

20

25

30

Met Gly Phe Leu Ser Leu Ile Ile Ser Pro Val Gly Met Val Leu Asn

35

40

45

Ser Thr Val Leu Trp Phe Leu Gly Phe Gln Ile Arg Arg Asn Ala Phe

50

55

60

Ser Val Tyr Ile Leu Asn Leu Ala Gly Ala Asp Phe Leu Phe Leu His

65

70

75

80

Ser Gln Phe Leu Phe Tyr Leu Leu Ala Ile Phe Pro Ser Ile Pro Ile

85

90

95

Gln Ile Pro Leu Phe Phe Asp Met Leu Thr Lys Phe Ala Tyr Leu Ser

100

105

110

Gly Leu Ser Ile Leu Ser Thr Ile Ser Ile Glu Arg Cys Leu Cys Val  
115 120 125

Met Trp Pro Ile Trp Tyr Arg Cys Gln Arg Pro Arg His Thr Ser Ser  
130 135 140

Val Thr Cys Ser Leu Leu Trp Ala Leu Ser Leu Leu Phe Ala Leu Leu  
145 150 155 160

Asp Gly Met Gly Cys Gly Leu Leu Phe Asn Ser Phe Asp Gln Ser Trp  
165 170 175

Cys Leu Lys Phe Asp Leu Ile Ile Cys Ala Trp Ser Ile Val Leu Phe  
180 185 190

Val Val Leu Cys Gly Ser Ser Leu Ile Leu Leu Val Arg Ile Phe Cys  
195 200 205

Gly Ser Gln Gln Ile Pro Val Thr Arg Leu Tyr Val Thr Ile Ala Leu  
210 215 220

Thr Val Leu Phe Phe Leu Ile Cys Cys Leu Pro Phe Gly Ile Ser Trp  
225 230 235 240

Ile Ile Gln Trp Ser Glu Thr Leu Ile Tyr Val Gly Phe Cys Asp Tyr  
245 250 255

Phe His Glu Glu Leu Phe Leu Ser Cys Ile Asn Ser Cys Ala Asn Pro  
260 265 270

Ile Ile Tyr Phe Leu Val Gly Phe Ile Arg Gln Arg Lys Phe Gln Gln  
275 280 285

Lys Ser Leu Lys Val Leu Leu Gln Arg Ala Met Glu Asp Thr Pro Glu  
290 295 300

Glu Glu Asn Glu Asp Met Gly Pro Ser Arg Asn Pro Glu Glu Phe Glu  
305 310 315 320

Thr Val Cys Ser Asn

325

<210> 4

<211> 330

<212> PRT

<213> Homo sapiens

<400> 4

Met Asp Pro Thr Thr Pro Ala Trp Gly Thr Glu Ser Thr Thr Val Asn  
1 5 10 15

Gly Asn Asp Gln Ala Leu Leu Leu Cys Gly Lys Glu Thr Leu Ile  
20 25 30

Pro Val Phe Leu Ile Leu Phe Ile Ala Leu Val Gly Leu Val Gly Asn  
35 40 45

Gly Phe Val Leu Trp Leu Leu Gly Phe Arg Met Arg Arg Asn Ala Phe  
50 55 60

Ser Val Tyr Val Leu Ser Leu Ala Gly Ala Asp Phe Leu Phe Leu Cys  
65 70 75 80

Phe Gln Ile Ile Asn Cys Leu Val Tyr Leu Ser Asn Phe Phe Cys Ser  
85 90 95

Ile Ser Ile Asn Phe Pro Ser Phe Phe Thr Thr Val Met Thr Cys Ala  
100 105 110

Tyr Leu Ala Gly Leu Ser Met Leu Ser Thr Val Ser Thr Glu Arg Cys  
115 120 125

Leu Ser Val Leu Trp Pro Ile Trp Tyr Arg Cys Arg Arg Pro Arg His  
130 135 140

Leu Ser Ala Val Val Cys Val Leu Leu Trp Ala Leu Ser Leu Leu Leu  
145 150 155 160

Ser Ile Leu Glu Gly Lys Phe Cys Gly Phe Leu Phe Ser Asp Gly Asp  
165 170 175

Ser Gly Trp Cys Gln Thr Phe Asp Phe Ile Thr Ala Ala Trp Leu Ile  
180 185 190

Phe Leu Phe Met Val Leu Cys Gly Ser Ser Leu Ala Leu Leu Val Arg  
195 200 205

Ile Leu Cys Gly Ser Arg Gly Leu Pro Leu Thr Arg Leu Tyr Leu Thr  
210 215 220

Ile Leu Leu Thr Val Leu Val Phe Leu Leu Cys Gly Leu Pro Phe Gly  
225 230 235 240

Ile Gln Trp Phe Leu Ile Leu Trp Ile Trp Lys Asp Ser Asp Val Leu  
245 250 255

Phe Cys His Ile His Pro Val Ser Val Val Leu Ser Ser Leu Asn Ser  
260 265 270

Ser Ala Asn Pro Ile Ile Tyr Phe Phe Val Gly Ser Phe Arg Lys Gln  
275 280 285

Trp Arg Leu Gln Gln Pro Ile Leu Lys Leu Ala Leu Gln Arg Ala Leu  
290 295 300

Gln Asp Ile Ala Glu Val Asp His Ser Glu Gly Cys Phe Arg Gln Gly  
305 310 315 320

Thr Pro Glu Met Ser Arg Ser Ser Leu Val

325

330

<210> 5

<211> 993

<212> DNA

<213> Mus musculus

<400> 5

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atcccaaaat tgatgatcat catcttcgga ctggtcggac tggatggaaa cgccattgtg 180  
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gccctggctg acttcctttt cctcctcagt agtacatag cttccaccct gtttcttctc 300  
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<210> 6

<211> 966

<212> DNA

<213> Homo sapiens

<400> 6

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gggatggcag gcaacagcat ggtgatctgg ctgctggct ttcaatgca caggaacccc 180  
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<210> 7

<211> 978

<212> DNA

<213> Mus musculus

<400> 7

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<210> 8

<211> 1770

<212> DNA

<213> Homo sapiens

<400> 8

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<211> 24

<212> DNA

<213> *Mus musculus*

<400> 9

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24

<210> 10

<211> 20

<212> DNA

<213> *Mus musculus*

<400> 10

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20

<210> 11

<211> 19

<212> DNA

<213> *Mus musculus*

<400> 11

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19

<210> 12  
<211> 20  
<212> DNA  
<213> Mus musculus

<400> 12  
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<210> 13  
<211> 21  
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<213> Homo sapiens

<400> 13  
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<210> 14  
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<212> DNA  
<213> Homo sapiens

<400> 14  
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<210> 15  
<211> 653  
<212> DNA  
<213> Mus musculus

<400> 15  
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cagtgctcac tccaaagcca cctctgaggt ccaggttagag gctttcata aaggctctgc 180  
tttgttactt gacatctcta ccatgttttc agctgtctca ggagtgtcct gcagtgcact 240  
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gtagatgatg gggttggcac agctgttaat agcagtcaagg acttctaattt ctagataaaag 360  
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aaagggcacc cgcagaggag aaaaacccaaa agggtcanca agatgggcac atgaaatctg 480  
gnaagcttta tntgccccag cgccacaaaaa acaacctggc canaaaaaac cngngntggn 540  
cangacnggg nncccnccc ccaaaanttt ttttntttn ctgnccnggg gnggnccctt 600  
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<210> 16

<211> 22

<212> DNA

<213> Homo sapiens

<400> 16

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22

<210> 17

<211> 21

<212> DNA

<213> Homo sapiens

<400> 17

ctaaggcgtt ggtgggtgcag g

21

<210> 18

<211> 25

<212> DNA

<213> Homo sapiens

<400> 18

TCTTAACCAC CAGATCATTC CTTCT

25

<210> 19  
<211> 20  
<212> DNA  
<213> Homo sapiens

<400> 19  
GGATACTGCG AGCAAATGGG 20

<210> 20  
<211> 24  
<212> DNA  
<213> Homo sapiens

<400> 20  
GGAGTCAGCC CTAAACTATT CCAG 24

<210> 21  
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